

SEQUENCE LISTING

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<120> CRYSTAL OF A GDP-FUCOSE SYNTHETASE POLYPEPTIDE

<130> W2025-701740

<140> 10/090,879
<141> 2002-03-04

<150> 09/373,432
<151> 1999-08-13

<150> 60/096,452
<151> 1998-08-13

<160> 3

<170> PatentIn version 3.5

<210> 1
<211> 338
<212> PRT
<213> Escherichia coli

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Cys Val Gln Leu Leu Gln Asn Gly His Asp Val Ile Ile Leu Asp Asn
20 25 30

Leu Cys Asn Ser Lys Arg Ser Val Leu Pro Val Ile Glu Arg Leu Gly
35 40 45

Gly Lys His Pro Thr Phe Val Glu Gly Asp Ile Arg Asn Glu Ala Leu
50 55 60

Met Thr Glu Ile Leu His Asp His Ala Ile Asp Thr Val Ile His Phe
65 70 75 80

Ala Gly Leu Lys Ala Val Gly Glu Ser Val Gln Lys Pro Leu Glu Tyr
85 90 95

Tyr Asp Asn Asn Val Asn Gly Thr Leu Arg Leu Ile Ser Ala Met Arg
100 105 110

Ala Ala Asn Val Lys Asn Phe Ile Phe Ser Ser Ser Ala Thr Val Tyr
 115 120 125

Gly Asp Asn Pro Lys Ile Pro Tyr Val Glu Ser Phe Pro Thr Gly Thr
 130 135 140

Pro Gln Ser Pro Tyr Gly Lys Ser Lys Leu Met Val Glu Gln Ile Leu
 145 150 155 160

Thr Asp Leu Gln Lys Ala Gln Pro Asp Trp Ser Ile Ala Leu Leu Arg
 165 170 175

Tyr Phe Asn Pro Val Gly Ala His Pro Ser Gly Asp Met Gly Glu Asp
 180 185 190

Pro Gln Gly Ile Pro Asn Asn Leu Met Pro Tyr Ile Ala Gln Val Ala
 195 200 205

Val Gly Arg Arg Asp Ser Leu Ala Ile Phe Gly Asn Asp Tyr Pro Thr
 210 215 220

Glu Asp Gly Thr Gly Val Arg Asp Tyr Ile His Val Met Asp Leu Ala
 225 230 235 240

Asp Gly His Val Val Ala Met Glu Lys Leu Ala Asn Lys Pro Gly Val
 245 250 255

His Ile Tyr Asn Leu Gly Ala Gly Val Gly Asn Ser Val Leu Asp Val
 260 265 270

Val Asn Ala Phe Ser Lys Ala Cys Gly Lys Pro Val Asn Tyr His Phe
 275 280 285

Ala Pro Arg Arg Glu Gly Asp Leu Pro Ala Tyr Trp Ala Asp Ala Ser
 290 295 300

Lys Ala Asp Arg Glu Leu Asn Trp Arg Val Thr Arg Thr Leu Asp Glu
 305 310 315 320

Met Ala Gln Asp Thr Trp His Trp Gln Ser Arg His Pro Gln Gly Tyr
 325 330 335

Pro Asp

<210> 2
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<212> PRT
<213> Escherichia coli

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1 5 10 15

Ile Arg Arg Gln Leu Glu Gln Arg Gly Asp Val Glu Leu Val Leu Arg
20 25 30

Thr Arg Asp Glu Leu Asn Leu Leu Asp Ser Arg Ala Val His Asp Phe
35 40 45

Phe Ala Ser Glu Arg Ile Asp Gln Val Tyr Leu Ala Ala Ala Lys Val
50 55 60

Gly Gly Ile Val Ala Asn Asn Thr Tyr Pro Ala Asp Phe Ile Tyr Gln
65 70 75 80

Asn Met Met Ile Glu Ser Asn Ile Ile His Ala Ala His Gln Asn Asp
85 90 95

Val Asn Lys Leu Leu Phe Leu Gly Ser Ser Cys Ile Tyr Pro Lys Leu
100 105 110

Ala Lys Gln Pro Met Ala Glu Ser Glu Leu Leu Gln Gly Thr Leu Glu
115 120 125

Pro Thr Asn Glu Pro Tyr Ala Ile Ala Lys Ile Ala Gly Ile Lys Leu
130 135 140

Cys Glu Ser Tyr Asn Arg Gln Tyr Gly Arg Asp Tyr Arg Ser Val Met
145 150 155 160

Pro Thr Asn Leu Tyr Gly Pro His Asp Asn Phe His Pro Ser Asn Ser
165 170 175

His Val Ile Pro Ala Leu Leu Arg Arg Phe His Glu Ala Thr Ala Gln
180 185 190

Asn Ala Pro Asp Val Val Val Trp Gly Ser Gly Thr Pro Met Arg Glu
195 200 205

Phe Leu His Val Asp Asp Met Ala Ala Ala Ser Ile His Val Met Glu
 210 215 220

Leu Ala His Glu Val Trp Leu Glu Asn Thr Gln Pro Met Leu Ser His
 225 230 235 240

Ile Asn Val Gly Thr Gly Val Asp Cys Thr Ile Arg Glu Leu Ala Gln
 245 250 255

Thr Ile Ala Lys Val Val Gly Tyr Lys Gly Arg Val Val Phe Asp Ala
 260 265 270

Ser Lys Pro Asp Gly Thr Pro Arg Lys Leu Leu Asp Val Thr Arg Leu
 275 280 285

His Gln Leu Gly Trp Tyr His Glu Ile Ser Leu Glu Ala Gly Leu Ala
 290 295 300

Ser Thr Tyr Gln Trp Phe Leu Glu Asn Gln Asp Arg Phe
 305 310 315

<210> 3
<211> 314
<212> PRT
<213> Homo sapiens

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Met Arg Ile Leu Val Thr Gly Gly Ser Gly Leu Val Gly Lys Ala Ile
 1 5 10 15

Gln Lys Val Val Ala Asp Gly Ala Gly Leu Pro Gly Glu Asp Trp Val
 20 25 30

Phe Val Ser Ser Lys Asp Ala Asp Leu Thr Asp Thr Ala Gln Thr Arg
 35 40 45

Ala Leu Pro Glu Lys Val Gln Pro Thr His Val Ile His Leu Ala Ala
 50 55 60

Met Val Gly Gly Leu Phe Arg Asn Ile Lys Tyr Asn Leu Asp Phe Trp
 65 70 75 80

Arg Lys Asn Val His Met Asn Asp Asn Val Leu His Ser Ala Phe Glu
 85 90 95

Val Gly Ala Arg Lys Val Val Ser Cys Leu Ser Thr Cys Ile Phe Pro
100 105 110

Asp Lys Thr Thr Tyr Pro Ile Asp Glu Thr Met Ile His Asn Gly Pro
115 120 125

Pro His Asn Ser Asn Phe Gly Tyr Ser Tyr Ala Lys Arg Met Ile Asp
130 135 140

Val Gln Asn Arg Ala Tyr Phe Gln Gln Tyr Gly Cys Thr Phe Thr Ala
145 150 155 160

Val Ile Pro Thr Asn Val Phe Gly Pro His Asp Asn Phe Asn Ile Glu
165 170 175

Asp Gly His Val Leu Pro Gly Leu Ile His Lys Val His Leu Ala Lys
180 185 190

Ser Ser Gly Ser Ala Leu Thr Val Trp Gly Thr Gly Asn Pro Arg Arg
195 200 205

Gln Phe Ile Tyr Ser Leu Asp Leu Ala Gln Leu Phe Ile Trp Val Leu
210 215 220

Arg Glu Tyr Asn Glu Val Glu Pro Ile Ile Leu Ser Val Gly Glu Glu
225 230 235 240

Asp Glu Val Ser Ile Lys Glu Ala Ala Glu Ala Val Val Glu Ala Met
245 250 255

Asp Phe His Gly Glu Val Thr Phe Asp Thr Thr Lys Ser Asp Gly Gln
260 265 270

Phe Lys Lys Thr Ala Ser Asn Ser Lys Leu Arg Thr Tyr Leu Pro Asp
275 280 285

Phe Arg Phe Thr Pro Phe Lys Gln Ala Val Lys Glu Thr Cys Ala Trp
290 295 300

Phe Thr Asp Asn Tyr Glu Gln Ala Arg Lys
305 310